

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/029,413

DATE: 01/15/2002
TIME: 07:55:35

Input Set : A:\ES.txt
Output Set: N:\CRF3\01152002\J029413.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: Malouf, Nadia
4 Nichols, Timothy
6 <120> TITLE OF INVENTION: Purified and Isolated Platelet Calcium Channel Nucleic Acids
and
7 Polypeptides and Therapeutic and Screening Methods Using Same
9 <130> FILE REFERENCE: 421/29
11 <140> CURRENT APPLICATION NUMBER: US/10/029,413
11 <141> CURRENT FILING DATE: 2001-12-20
11 <160> NUMBER OF SEQ ID NOS: 29
13 <170> SOFTWARE: PatentIn version 3.0

ERRORED SEQUENCES

2714 <210> SEQ ID NO: 10
2715 <211> LENGTH: 1338
2716 <212> TYPE: PRT
2717 <213> ORGANISM: Homo sapiens
2719 <400> SEQUENCE: 10
2721 Met Ser Lys Gly Leu Pro Asp Lys Ser Glu Glu Glu Lys Ser Thr Met
2722 1 5 10 15
2725 Ala Lys Lys Leu Glu Gln Lys Pro Lys Gly Glu Gly Ile Pro Thr Thr
2726 20 25 30
E--> 2729 Ala Lys Leu Xaa Ile Asp Glu Xaa Glu Ser Asn Val Asn Glu Val Lys
2730 35 40 45
2733 Asp Pro Tyr Pro Ser Ala Asp Phe Pro Gly Asp Asp Glu Glu Asp Glu
2734 50 55 60
2737 Pro Glu Ile Pro Leu Ser Pro Arg Pro Arg Pro Leu Ala Glu Leu Gln
2738 65 70 75 80
2741 Leu Lys Glu Lys Ala Val Pro Ile Pro Glu Ala Ser Ser Phe Phe Ile
2742 85 90 95
2745 Phe Ser Pro Thr Asn Lys Ile Arg Val Leu Cys His Arg Ile Val Asn
2746 100 105 110
2749 Ala Thr Trp Phe Thr Asn Phe Ile Leu Leu Phe Ile Leu Leu Ser Ser
2750 115 120 125
2753 Ala Ala Leu Ala Ala Glu Asp Pro Ile Arg Ala Asp Ser Met Arg Asn
2754 130 135 140
2757 Gln Ile Leu Lys His Phe Asp Ile Gly Phe Thr Ser Val Phe Thr Val
2758 145 150 155 160
2761 Glu Ile Val Leu Lys Met Thr Thr Tyr Gly Ala Phe Leu His Lys Gly
2762 165 170 175
2765 Ser Phe Cys Arg Asn Tyr Phe Asn Met Leu Asp Leu Leu Val Val Ala
2766 180 185 190
2769 Val Ser Leu Ile Ser Met Gly Leu Glu Ser Ser Ala Ile Ser Val Val
2770 195 200 205
2773 Lys Ile Leu Arg Val Leu Arg Val Leu Arg Pro Leu Arg Ala Ile Asn
2774 210 215 220
E--> 2777 Arg Ala Lys Gly Leu Lys Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa

pp 1-2,4

*see item 9
on Enov
Summary*

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Input Set : A:\ES.txt
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2778 225 230 235 240
E--> 2781 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
2782 245 250 255
E--> 2785 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
2786 260 265 270
E--> 2789 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
2790 275 280 285
E--> 2793 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
2794 290 295 300
E--> 2797 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
2798 305 310 315 320
E--> 2801 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys His Arg Gly Tyr
2802 325 330 335
2805 Tyr Tyr Val Tyr Lys Asp Gly Asp Pro Met Gln Ile Glu Leu Arg His
2806 340 345 350
2809 Arg Glu Trp Val His Ser Asp Phe His Phe Asp Asn Val Leu Ser Ala
2810 355 360 365
2813 Met Met Ser Leu Phe Thr Val Ser Thr Phe Glu Gly Trp Pro Gln Leu
2814 370 375 380
2817 Leu Tyr Lys Ala Ile Asp Ser Asn Ala Glu Asp Val Gly Pro Ile Tyr
2818 385 390 395 400
2821 Asn Asn Arg Val Glu Met Ala Ile Phe Phe Ile Ile Tyr Ile Ile Leu
2822 405 410 415
2825 Ile Ala Phe Phe Met Met Asn Ile Phe Val Gly Phe Val Ile Val Thr
2826 420 425 430
2829 Phe Gln Glu Gln Gly Glu Thr Glu Tyr Lys Asn Cys Glu Leu Asp Lys
2830 435 440 445
2833 Asn Gln Arg Gln Cys Val Gln Tyr Ala Leu Lys Ala Arg Pro Leu Arg
2834 450 455 460
2837 Cys Tyr Ile Pro Lys Asn Pro Tyr Gln Tyr Gln Val Trp Tyr Ile Val
2838 465 470 475 480
2841 Thr Ser Ser Tyr Phe Glu Tyr Leu Met Phe Ala Leu Ile Met Leu Asn
2842 485 490 495
2845 Thr Ile Cys Leu Gly Met Gln His Tyr Asn Gln Ser Glu Gln Met Asn
2846 500 505 510
2849 His Ile Ser Asp Ile Leu Asn Val Ala Phe Thr Ile Ile Phe Thr Leu
2850 515 520 525
2853 Glu Met Ile Leu Lys Leu Met Ala Phe Lys Ala Arg Gly Tyr Phe Gly
2854 530 535 540
2857 Asp Pro Trp Asn Val Phe Asp Phe Leu Ile Val Ile Gly Ser Ile Ile
2858 545 550 555 560
2861 Asp Val Ile Leu Ser Glu Ile Asp Thr Phe Leu Ala Ser Ser Gly Gly
2862 565 570 575
2865 Leu Tyr Cys Leu Gly Gly Gly Cys Gly Asn Val Asp Pro Asp Glu Ser
2866 580 585 590
2869 Ala Arg Ile Ser Ser Ala Phe Phe Arg Leu Phe Arg Val Met Arg Leu
2870 595 600 605
2873 Ile Lys Leu Leu Ser Arg Ala Glu Gly Val Arg Thr Leu Leu Trp Thr
2874 610 615 620

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2877 Phe Ile Lys Ser Phe Gln Ala Leu Pro Tyr Val Ala Leu Leu Ile Val
2878 625 630 635 640
2881 Met Leu Phe Phe Ile Tyr Ala Val Ile Gly Met Gln Met Phe Gly Lys
2882 645 650 655
2885 Ile Ala Leu Val Asp Gly Thr Gln Ile Asn Arg Asn Asn Asn Phe Gln
2886 660 665 670
2889 Thr Phe Pro Gln Ala Val Leu Leu Leu Phe Arg Cys Ala Thr Gly Glu
2890 675 680 685
2893 Ala Trp Gln Glu Ile Leu Leu Ala Cys Ser Tyr Gly Lys Leu Cys Asp
2894 690 695 700
2897 Pro Glu Ser Asp Tyr Ala Pro Gly Glu Glu Tyr Thr Cys Gly Thr Asn
2898 705 710 715 720
2901 Phe Ala Tyr Tyr Tyr Phe Ile Ser Phe Tyr Met Leu Cys Ala Phe Leu
2902 725 730 735
2905 Val Ile Asn Leu Phe Val Ala Val Ile Met Asp Asn Phe Asp Tyr Leu
2906 740 745 750
2909 Thr Arg Asp Trp Ser Ile Leu Gly Pro His His Leu Asp Glu Phe Lys
2910 755 760 765
2913 Ala Ile Trp Ala Glu Tyr Asp Pro Glu Ala Lys Gly Arg Ile Lys His
2914 770 775 780
2917 Leu Asp Val Val Thr Leu Leu Arg Arg Ile Gln Pro Pro Leu Gly Phe
2918 785 790 795 800
2921 Gly Lys Phe Cys Pro His Arg Val Ala Cys Lys Arg Leu Val Gly Met
2922 805 810 815
2925 Asn Met Pro Leu Asn Ser Asp Gly Thr Val Thr Phe Asn Ala Thr Leu
2926 820 825 830
2929 Phe Ala Leu Val Arg Thr Ala Leu Lys Ile Lys Thr Glu Gly Asn Phe
2930 835 840 845
2933 Glu Gln Ala Asn Glu Glu Leu Arg Ala Ile Ile Lys Lys Ile Trp Lys
2934 850 855 860
2937 Arg Thr Ser Met Lys Leu Leu Asp Gln Val Ile Pro Pro Ile Gly Asp
2938 865 870 875 880
2941 Asp Glu Val Thr Val Gly Lys Phe Tyr Ala Thr Phe Leu Ile Gln Glu
2942 885 890 895
2945 His Phe Arg Lys Phe Met Lys Arg Gln Glu Glu Tyr Tyr Gly Tyr Arg
2946 900 905 910
2949 Pro Lys Lys Asp Ile Val Gln Ile Gln Ala Gly Leu Arg Thr Ile Glu
2950 915 920 925
2953 Glu Glu Ala Ala Pro Glu Ile Cys Arg Thr Val Ser Gly Asp Leu Ala
2954 930 935 940
2957 Ala Glu Glu Glu Leu Glu Arg Ala Met Val Glu Ala Ala Met Glu Glu
2958 945 950 955 960
2961 Gly Ile Phe Arg Arg Thr Gly Gly Leu Phe Gly Gln Val Asp Asn Phe
2962 965 970 975
2965 Leu Glu Arg Thr Asn Ser Leu Pro Pro Val Met Ala Asn Gln Arg Pro
2966 980 985 990
2969 Leu Gln Phe Ala Glu Ile Glu Met Glu Glu Met Glu Ser Pro Val Phe
2970 995 1000 1005
2973 Leu Glu Asp Phe Pro Gln Asp Pro Arg Thr Asn Pro Leu Ala Arg

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2974	1010	1015	1020
2977	Ala Asn Thr Asn Asn Ala Asn Ala Asn Val Ala Tyr Gly Asn Ser		
2978	1025	1030	1035
2981	Asn His Ser Asn Ser His Val Phe Ser Ser Val His Tyr Glu Arg		
2982	1040	1045	1050
2985	Glu Phe Pro Glu Glu Thr Glu Thr Pro Ala Thr Arg Gly Arg Ala		
2986	1055	1060	1065
E--> 2989	Leu Gly Gln Pro Cys Arg Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa		
2990	1070	1075	1080
E--> 2993	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa		
2994	1085	1090	1095
E--> 2997	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa		
2998	1100	1105	1110
E--> 3001	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa		
3002	1115	1120	1125
E--> 3005	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa		
3006	1130	1135	1140
E--> 3009	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa		
3010	1145	1150	1155
E--> 3013	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa		
3014	1160	1165	1170
E--> 3017	Xaa Xaa Thr Gly Pro His Ser Lys Pro Cys Val Glu Met Leu Lys		
3018	1175	1180	1185
3021	Gly Leu Leu Thr Gln Arg Ala Met Pro Arg Gly Gln Ala Pro Pro		
3022	1190	1195	1200
3025	Ala Pro Cys Gln Cys Pro Arg Val Glu Ser Ser Met Pro Glu Asp		
3026	1205	1210	1215
3029	Arg Lys Ser Ser Thr Pro Gly Ser Leu His Glu Glu Thr Pro His		
3030	1220	1225	1230
3033	Ser Arg Ser Thr Arg Glu Asn Thr Ser Arg Cys Ser Ala Pro Ala		
3034	1235	1240	1245
3037	Thr Ala Leu Leu Ile Gln Lys Ala Leu Val Arg Gly Gly Leu Gly		
3038	1250	1255	1260
3041	Thr Leu Ala Ala Asp Ala Asn Phe Ile Met Ala Thr Gly Gln Ala		
3042	1265	1270	1275
3045	Leu Ala Asp Ala Cys Gln Met Glu Pro Glu Glu Val Glu Ile Met		
3046	1280	1285	1290
3049	Ala Thr Glu Leu Leu Lys Gly Arg Glu Ala Pro Glu Gly Met Ala		
3050	1295	1300	1305
3053	Ser Ser Leu Gly Cys Leu Asn Leu Gly Ser Ser Leu Gly Ser Leu		
3054	1310	1315	1320
3057	Asp Gln His Gln Gly Ser Gln Glu Thr Leu Ile Pro Pro Arg Leu		
3058	1325	1330	1335
5629	<210> SEQ ID NO: 16		
5630	<211> LENGTH: 1772		
5631	<212> TYPE: PRT		
5632	<213> ORGANISM: Homo sapiens		
5634	<400> SEQUENCE: 16		
5636	Met Pro Thr Ser Glu Thr Glu Ser Val Asn Thr Glu Asn Val Ser Gly		

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```

5637 1          5          10          15
5640 Glu Gly Glu Asn Arg Gly Cys Cys Gly Ser Leu Trp Cys Trp Trp Arg
5641          20          25          30
5644 Arg Arg Gly Ala Ala Lys Ala Gly Pro Ser Gly Cys Arg Arg Trp Gly
5645          35          40          45
5648 Gln Ala Ile Ser Lys Ser Lys Leu Ser Arg Arg Trp Arg Arg Trp Asn
5649          50          55          60
5652 Arg Phe Asn Arg Arg Arg Cys Arg Ala Ala Val Lys Ser Val Thr Phe
5653 65          70          75          80
5656 Tyr Trp Leu Val Ile Val Leu Val Phe Leu Asn Thr Leu Thr Ile Ser
5657          85          90          95
5660 Ser Glu His Tyr Asn Gln Pro Asp Trp Leu Thr Gln Ile Gln Asp Met
5661          100          105          110
5664 Pro Asn Lys Val Leu Leu Ala Leu Phe Thr Cys Glu Met Leu Val Lys
5665          115          120          125
5668 Met Tyr Ser Leu Gly Leu Gln Ala Tyr Phe Val Ser Leu Phe Asn Arg
5669          130          135          140
5672 Phe Asp Cys Phe Val Val Cys Gly Gly Ile Thr Glu Thr Ile Leu Val
5673 145          150          155          160
5676 Glu Leu Glu Ile Met Ser Pro Leu Gly Ile Ser Val Phe Arg Cys Val
5677          165          170          175
5680 Arg Leu Leu Arg Ile Phe Lys Val Thr Arg His Trp Thr Ser Leu Ser
5681          180          185          190
5684 Asn Leu Val Ala Ser Leu Leu Asn Ser Met Lys Ser Ile Ala Ser Leu
5685          195          200          205
5688 Leu Leu Leu Leu Phe Leu Phe Ile Ile Ile Phe Ser Leu Leu Gly Met
5689          210          215          220
5692 Gln Leu Phe Gly Gly Lys Phe Asn Phe Asp Glu Thr Gln Thr Lys Arg
5693 225          230          235          240
5696 Ser Thr Phe Asp Asn Phe Pro Gln Ala Leu Leu Thr Val Phe Gln Ile
5697          245          250          255
5700 Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr Asp Gly Ile Met Ala
5701          260          265          270
5704 Tyr Gly Gly Pro Ser Ser Ser Gly Met Ile Val Cys Ile Tyr Phe Ile
5705          275          280          285
5708 Ile Leu Phe Ile Cys Gly Asn Tyr Ile Leu Leu Asn Val Phe Leu Ala
5709          290          295          300
5712 Ile Ala Val Asp Asn Leu Ala Asp Ala Glu Ser Leu Asn Thr Ala Gln
5713 305          310          315          320
5716 Lys Glu Glu Ala Glu Glu Lys Glu Arg Lys Lys Ile Ala Arg Lys Glu
5717          325          330          335
5720 Ser Leu Glu Asn Lys Lys Asn Asn Lys Pro Glu Val Asn Gln Ile Ala
5721          340          345          350
5724 Asn Ser Asp Asn Lys Val Thr Ile Asp Asp Tyr Arg Glu Glu Asp Glu
5725          355          360          365
5728 Asp Lys Asp Pro Tyr Pro Pro Cys Asp Val Pro Gly Met Val Xaa Xaa
5729          370          375          380
E--> 5732 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
5733 385          390          395          400

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5833 785          790          795          800
5836 Pro Lys Asn Pro Tyr Gln Tyr Lys Phe Trp Tyr Val Val Asn Ser Ser
5837          805          810          815
5840 Pro Phe Glu Tyr Met Met Phe Val Leu Ile Met Leu Asn Thr Leu Cys
5841          820          825          830
5844 Leu Ala Met Gln His Tyr Glu Gln Ser Lys Met Phe Asn Asp Ala Met
5845          835          840          845
5848 Asp Ile Leu Asn Met Val Phe Thr Gly Val Phe Thr Val Glu Met Val
5849          850          855          860
5852 Leu Lys Val Ile Ala Phe Lys Pro Lys Gly Tyr Phe Ser Asp Ala Trp
5853 865          870          875          880
5856 Asn Thr Phe Asp Ser Leu Ile Val Ile Gly Ser Ile Ile Asp Val Ala
5857          885          890          895
5860 Leu Ser Glu Ala Lys Pro Thr Glu Ser Glu Asn Val Pro Val Pro Thr
5861          900          905          910
5864 Ala Thr Pro Gly Asn Ser Glu Glu Ser Asn Arg Ile Ser Ile Thr Phe
5865          915          920          925
5868 Phe Arg Leu Phe Arg Val Met Arg Leu Val Lys Leu Leu Ser Arg Gly
5869          930          935          940
5872 Glu Gly Ile Arg Thr Leu Leu Trp Thr Phe Ile Lys Ser Phe Gln Ala
5873 945          950          955          960
5876 Leu Pro Tyr Val Ala Leu Leu Ile Ala Met Leu Phe Phe Ile Tyr Ala
5877          965          970          975
5880 Val Ile Gly Met Gln Met Phe Gly Lys Val Ala Met Arg Asp Asn Asn
5881          980          985          990
5884 Gln Ile Asn Arg Asn Asn Asn Phe Gln Thr Phe Pro Gln Ala Val Leu
5885          995          1000          1005
5888 Leu Leu Phe Arg Cys Ala Thr Gly Glu Ala Trp Gln Glu Ile Met
5889          1010          1015          1020
5892 Leu Ala Cys Leu Pro Gly Lys Leu Cys Asp Pro Glu Ser Asp Tyr
5893          1025          1030          1035
5896 Asn Pro Gly Glu Glu Tyr Thr Cys Gly Ser Asn Phe Ala Ile Val
5897          1040          1045          1050
5900 Tyr Phe Ile Ser Phe Tyr Met Leu Cys Ala Phe Leu Ile Ile Asn
5901          1055          1060          1065
5904 Leu Phe Val Ala Val Ile Met Asp Asn Phe Asp Tyr Leu Thr Arg
5905          1070          1075          1080
5908 Asp Trp Ser Ile Leu Gly Pro His His Leu Asp Glu Phe Lys Arg
5909          1085          1090          1095
5912 Ile Trp Ser Glu Tyr Asp Pro Glu Ala Lys Gly Arg Ile Lys His
5913          1100          1105          1110
5916 Leu Asp Val Val Thr Leu Leu Arg Arg Ile Gln Pro Pro Leu Gly
5917          1115          1120          1125
5920 Phe Gly Lys Leu Cys Pro His Arg Val Ala Cys Lys Arg Leu Val
5921          1130          1135          1140
5924 Ala Met Asn Met Pro Leu Asn Ser Asp Gly Thr Val Met Phe Asn
5925          1145          1150          1155
5928 Ala Thr Leu Phe Ala Leu Val Arg Thr Ala Leu Lys Ile Lys Thr
5929          1160          1165          1170

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5932	Glu	Gly	Asn	Leu	Glu	Gln	Ala	Asn	Glu	Glu	Leu	Arg	Ala	Val	Ile
5933		1175					1180					1185			
5936	Lys	Lys	Ile	Trp	Lys	Lys	Thr	Ser	Met	Lys	Leu	Leu	Asp	Gln	Val
5937		1190					1195					1200			
5940	Val	Pro	Pro	Ala	Gly	Asp	Asp	Glu	Val	Thr	Val	Gly	Lys	Phe	Tyr
5941		1205					1210					1215			
5944	Ala	Thr	Phe	Leu	Ile	Gln	Asp	Tyr	Phe	Arg	Lys	Phe	Lys	Lys	Arg
5945		1220					1225					1230			
5948	Lys	Glu	Gln	Gly	Leu	Val	Gly	Lys	Tyr	Pro	Ala	Lys	Asn	Thr	Thr
5949		1235					1240					1245			
5952	Ile	Ala	Leu	Gln	Ala	Gly	Leu	Arg	Thr	Leu	His	Asp	Ile	Gly	Pro
5953		1250					1255					1260			
5956	Glu	Ile	Arg	Arg	Ala	Ile	Ser	Cys	Asp	Leu	Gln	Asp	Asp	Glu	Pro
5957		1265					1270					1275			
5960	Glu	Glu	Thr	Lys	Arg	Glu	Glu	Glu	Asp	Asp	Val	Phe	Lys	Arg	Asn
5961		1280					1285					1290			
5964	Gly	Ala	Leu	Leu	Gly	Asn	His	Val	Asn	His	Val	Asn	Ser	Asp	Arg
5965		1295					1300					1305			
5968	Arg	Asp	Ser	Leu	Gln	Gln	Thr	Asn	Thr	Thr	His	Arg	Pro	Leu	His
5969		1310					1315					1320			
5972	Val	Gln	Arg	Pro	Ser	Ile	Pro	Pro	Ala	Ser	Asp	Thr	Glu	Lys	Pro
5973		1325					1330					1335			
5976	Leu	Phe	Pro	Pro	Ala	Gly	Asn	Ser	Val	Cys	His	Asn	His	His	Asn
5977		1340					1345					1350			
5980	His	Asn	Ser	Ile	Gly	Lys	Gln	Val	Pro	Thr	Ser	Thr	Asn	Ala	Asn
5981		1355					1360					1365			
5984	Leu	Asn	Asn	Ala	Asn	Met	Ser	Lys	Ala	Ala	His	Gly	Lys	Arg	Pro
5985		1370					1375					1380			
5988	Ser	Ile	Gly	Asn	Leu	Glu	His	Val	Ser	Glu	Asn	Gly	His	His	Ser
5989		1385					1390					1395			
5992	Ser	His	Lys	His	Asp	Arg	Glu	Pro	Gln	Arg	Arg	Ser	Ser	Val	Lys
5993		1400					1405					1410			
5996	Arg	Thr	Arg	Tyr	Tyr	Glu	Thr	Tyr	Ile	Arg	Ser	Asp	Ser	Gly	Asp
5997		1415					1420					1425			
6000	Glu	Gln	Leu	Pro	Thr	Ile	Cys	Arg	Glu	Asp	Pro	Glu	Ile	His	Gly
6001		1430					1435					1440			
6004	Tyr	Phe	Arg	Asp	Pro	His	Cys	Leu	Gly	Glu	Gln	Glu	Tyr	Phe	Ser
6005		1445					1450					1455			
6008	Ser	Glu	Glu	Cys	Tyr	Glu	Asp	Asp	Ser	Ser	Pro	Thr	Trp	Ser	Arg
6009		1460					1465					1470			
6012	Gln	Asn	Tyr	Gly	Tyr	Tyr	Ser	Arg	Tyr	Pro	Gly	Arg	Asn	Ile	Asp
6013		1475					1480					1485			
6016	Ser	Glu	Arg	Pro	Arg	Gly	Tyr	His	His	Pro	Gln	Gly	Phe	Leu	Glu
6017		1490					1495					1500			
6020	Asp	Asp	Asp	Ser	Pro	Val	Cys	Tyr	Asp	Ser	Arg	Arg	Ser	Pro	Arg
6021		1505					1510					1515			
6024	Arg	Arg	Leu	Leu	Pro	Pro	Thr	Pro	Ala	Ser	His	Arg	Arg	Ser	Ser
6025		1520					1525					1530			
6028	Phe	Asn	Phe	Glu	Cys	Leu	Arg	Arg	Gln	Ser	Ser	Gln	Glu	Glu	Val

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6029	1535	1540	1545
6032	Pro Ser Ser Pro Ile Phe	Pro His Arg Thr Ala	Leu Pro Leu His
6033	1550	1555	1560
6036	Leu Met Gln Gln Gln Ile	Met Ala Val Ala Gly	Leu Asp Ser Ser
6037	1565	1570	1575
6040	Lys Ala Gln Lys Tyr Ser	Pro Ser His Ser Thr	Arg Ser Trp Ala
6041	1580	1585	1590
6044	Thr Pro Pro Ala Thr Pro	Pro Tyr Arg Asp Trp	Thr Pro Cys Tyr
6045	1595	1600	1605
6048	Thr Pro Leu Ile Gln Val	Glu Gln Ser Glu Ala	Leu Asp Gln Val
6049	1610	1615	1620
6052	Asn Gly Ser Leu Pro Ser	Leu His Arg Ser Ser	Trp Tyr Thr Asp
6053	1625	1630	1635
6056	Glu Pro Asp Ile Ser Tyr	Arg Thr Phe Thr Pro	Ala Ser Leu Thr
6057	1640	1645	1650
6060	Val Pro Ser Ser Phe Arg	Asn Lys Asn Ser Asp	Lys Gln Arg Ser
6061	1655	1660	1665
6064	Ala Asp Ser Leu Val Glu	Ala Val Leu Ile Ser	Glu Gly Leu Gly
6065	1670	1675	1680
6068	Arg Tyr Ala Arg Asp Pro	Lys Phe Val Ser Ala	Thr Lys His Glu
6069	1685	1690	1695
6072	Ile Ala Asp Ala Cys Asp	Leu Thr Ile Asp Glu	Met Glu Ser Ala
6073	1700	1705	1710
6076	Ala Ser Thr Leu Leu Asn	Gly Asn Val Arg Pro	Arg Ala Asn Gly
6077	1715	1720	1725
6080	Asp Val Gly Pro Leu Ser	His Arg Gln Asp Tyr	Glu Leu Gln Asp
6081	1730	1735	1740
6084	Phe Gly Pro Gly Tyr Ser	Asp Glu Glu Pro Asp	Pro Gly Arg Asp
6085	1745	1750	1755
6088	Glu Glu Asp Leu Ala Asp	Glu Met Ile Cys Ile	Thr Thr Leu
6089	1760	1765	1770

VERIFICATION SUMMARY

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Input Set : A:\ES.txt

Output Set: N:\CRF3\01152002\J029413.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:2145 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:2176 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6
L:2196 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:2200 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:2209 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:7
L:2238 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:8
L:2263 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:2369 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2370 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2417 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2418 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2421 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2422 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2425 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2426 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2429 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2430 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2433 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2434 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2437 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2438 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2441 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2442 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2629 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2630 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2633 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2634 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2637 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2638 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2641 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2642 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2645 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2646 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2649 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2650 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2653 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2654 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2657 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2658 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2729 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:10
M:340 Repeated in SeqNo=10
L:5246 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:5247 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:5250 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:5251 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:5254 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/029,413

DATE: 01/15/2002

TIME: 07:55:38

Input Set : A:\ES.txt

Output Set: N:\CRF3\01152002\J029413.raw

L:5255 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:5258 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:5259 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:5262 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:5263 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:5266 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:5267 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:5270 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:5271 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:5728 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:16
M:340 Repeated in SeqNo=16
L:9466 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:23
L:9496 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:25